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OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/824,735

DATE: 09/07/2001  
TIME: 14:11:35

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3 <110> APPLICANT: ZHU, JIAN-KANG  
4 LIU, JIPING  
5 ISHITANI, MANABU  
6 HALFTER, URSULA  
7 KIM, CHEOL-SOO  
9 <120> TITLE OF INVENTION: PROTEINS AND DNA RELATED TO SALT TOLERANCE IN PLANTS  
11 <130> FILE REFERENCE: 205645US20  
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C--> 14 <141> CURRENT FILING DATE: 2001-08-17  
16 <150> PRIOR APPLICATION NUMBER: US 60/824,735  
17 <151> PRIOR FILING DATE: 2000-04-04  
19 <160> NUMBER OF SEQ ID NOS: 16  
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179 15 20 25
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182 Phe Ala Arg Asn Thr Asp Thr Gly Asp Asn Val Ala Ile Lys Ile Met
183 30 35 40
185 gct aag agt aca ata ctt aag aac aga atg gtt gat cag gtatgttctg 2178
186 Ala Lys Ser Thr Ile Leu Lys Asn Arg Met Val Asp Gln
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191 taaagctctt ttacag ata aaa aga gag ata tct ata atg aag att gtt cgt 2290
192 Ile Lys Arg Glu Ile Ser Ile Met Lys Ile Val Arg
193 60 65
195 cac ccg aac ata gtg agg ttg tat gag gtatgtttgt ttgtttccat 2337
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197 70 75
199 gcatctgcga aattttatct ctgaagtgtt tttgcatcat tgttcttctg ttgttttttt 2397
201 gtgattttcc cgatgtag gtg ttg gcg agt cct tcg aaa ata tat ata gtt 2448
202 Val Leu Ala Ser Pro Ser Lys Ile Tyr Ile Val
203 80 85
205 ttg gag ttt gtg aca gga gga gag ctc ttt gat aga att gtacggaact 2497
206 Leu Glu Phe Val Thr Gly Gly Glu Leu Phe Asp Arg Ile
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209 tccatacttg taggcagcgt ccattagtta aaacctctct acttaatttt ttaatatatg 2557
211 aaatctttca tgcag gtt cat aaa ggg agg ctt gaa gaa agt gag tct cgg 2608
212 Val His Lys Gly Arg Leu Glu Glu Ser Glu Ser Arg
213 105 110
215 aaa tac ttt caa cag ctt gta gat gct gtt gct cat tgt cac tgc aag 2656
216 Lys Tyr Phe Gln Gln Leu Val Asp Ala Val Ala His Cys His Cys Lys
217 115 120 125
219 ggt gtt tac cac cgt gac cta aag gtaaagacgt gtttttgttt accaatattc 2710
220 Gly Val Tyr His Arg Asp Leu Lys
221 130 135
223 ctcaagaatat ctcaactgcgt tgcaatccag acttgatatt tttgtgtcgc tatgttatgt 2770
225 tatctag cca gaa aat ctt tta ctc gat aca aat gga aat ctg aag gtt 2819
226 Pro Glu Asn Leu Leu Leu Asp Thr Asn Gly Asn Leu Lys Val
227 140 145 150
229 tcg gat ttc gga ctc agt gca ttg cct cag gaa gtaagtgtc ttatctctgc 2872
230 Ser Asp Phe Gly Leu Ser Ala Leu Pro Gln Glu

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239 165          170          175          180
241 ctt agt gga cag ggt tac gat ggt tca gca gct gat att tgg tct tgc 3025
242 Leu Ser Gly Gln Gly Tyr Asp Gly Ser Ala Ala Asp Ile Trp Ser Cys
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245 ggg gtt att ctt ttc gtt ata ttg gct gga tat tta cct ttt tcc gag 3073
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270          Arg Ile Gln Ile Gln Gly Ile Lys Lys Asp
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275          265          270          275
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298                                     Gly Ser Tyr Val Ala Glu Asn
299          295
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VERIFICATION SUMMARY

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